FIGURE 2 A

TGTTATTGC TCTAGAGTC ATGAAACAAC AAAAACGGCT TTACGCCCGA TTGCTGACGC HgaI PstI XbaI

AATTICAGGA GAACACCTAA TTAAAGCCTG GACACATTIT CATITICCGG GGCGCGGCAG CACATACAGC GATTITAAAT GGCATTGGTA CCATTTTGAC GGAACCGATT GGGACGAGTC CCGAAAGCTG AACCGCATCT ATAAGTTTCA AGGAAAGGCT TGGGATTGGG AAGTTTCCAA CAGGGAAAAA ACGGGGAAGG AAATGTTTAC GGTAGCTGAA TATTGGCAGA ATGACTTGGG GCAGTATTTT GAATGGTACA TGCCCAATGA CGGCCAACAT TGGAAGCGTT TGCAAAACGA CTCGGCATAT TTGGCTGAAC ACGGTATTAC TGCCGTCTGG ATTCCCCCGG CATATAAGGG AACGAGCCAA GCGGATGTGG GCTACGGTGC TTACGACCTT TATGATTTAG GGGAGTTTCA TCAAAAAGGG ACGGTTCGGA CAAAGTACGG CACAAAAGGA GAGCTGCAAT CTGCGATCAA AAGTCTTCAT TCCCGCGACA TTAACGTTTA CGGGGATGTG GTCATCAACC ACAAAGGCGG CGCTGATGCG ACCGAAGATG TAACCGCGGT TGAAGTCGAT CCCGCTGACC GCAACCGCGT TGAAAACGGC AACTATGATT ATTTGATGTA TGCCGACATC GATTATGACC ATCCTGATGT CGCAGCAGAA ATTAAGAGAT GGGGCACTTG GTATGCCAAT GAACTGCAAT TGGACGGTTT CCGTCTTGAT GCTGTCAAAC ACATTAAATT TTCTTTTTG CGGGATTGGG TTAATCATGT GCTCATCTTC TTGCTGCCTC ATTCTGCAGC AGCGGCGGCA AATCTTAATG GGACGCTGAT

FIGURE 2B

GAACGGTACG GTCGTTTCCA AGCATCCGTT GAAATCGGTT ACATTTGTCG ATAACCATGA CGCGCTGGAA AACTATTTGA ACAAAACAAA TTTTAATCAT TCAGTGTTTG ACGTGCCGCT

ATGGCATGAC ATTACCGGAA ACCGTTCGGA GCCGGTTGTC ATCAATTCGG AAGGCTGGGG AACAGACGGA CCCGGTGGGG CAAAGCGAAT GTATGTCGGC CGGCAAAACG CCGGTGAGAC AGAGTITCAC GTAAACGGCG GGTCGGTTTC AATTTATGTT CAAAGATAGA AGAGCAGAGA CGCTTTTATT CTCACAAGGG AATCTGGATA CCCTCAGGTT TTCTACGGGG ATATGTACGG AAAAGCGAGA AAACAGTATG CGTACGGAGC ACAGCATGAT TATTTCGACC ACCATGACAT TGTCGGCTGG ACAAGGGAAG GCGACAGCTC GGTTGCAAAT TCAGGTTTGG CGGCATTAAT GACGAAAGGA GACTCCCAGC GCGAAATTCC TGCCTTGAAA CACAAAATTG AACCGATCTT TACACAGCCG GGGCAATCGC TTGAGTCGAC TGTCCAAACA TGGTTTAAGC CGCTTGCTTA

BamHI

GGACGGATTT CCTGAAGGAA ATCCGTTTTT TTATTTTGCC CGTCTTATAA ATTTCTTTGA TTACATTTTA TAATTATTT TAACAAAGTG TCATCAGCCC TCAGGAAGGA CTTGCTGACA GTTTGAATCG CATAGGTAAG GCGGGGATGA AATGGCAACG TTATCTGATG TAGCAAAGAA

AGCANATGTG TCGAAAATGA CGGTATCGCG GGTGATCA SEQ ID N

FIGURE 3

Oligonucleotide duplex A

	SEQ ID NO:6 SEQ ID NO:7			
NCOI BamHI H1DGIII	5' GGGTTTTTATTTTAATTTTCTTTCAAATACTTCCACCATGGGTAACGGATCCA 3' S	Oligonucleotide duplex B	$\frac{\text{HGaI SITE}}{\alpha-\text{AMYLASE}}$	catg gcaaatcttaatggacgctgatg 3' SEQ ID NO:8 cgtttagaattacctgcgactacgtcat 5' SEQ ID NO:9 Met mature α-Amylase

FIGURE 2 A

	TGTTATTTGC <u>Hqa</u> I
Haal	TCTAGAGTC ATGAAACAAC AAAAACGGCT TTACGCCCGA TTGCTGACGC TGTTATTTGC HGAI
	TTACGCCCGA
	AAAAACGGCT Psti
*	ATGAAACAAC
XbaI	TCTAGAGTC

CAGGGAAAAA ACGGGGAAGG AAATGTTTAC GGTAGCTGAA TATTGGCAGA ATGACTTGGG CGCAGCAGAA ATTAAGAGAT GGGGCACTTG GTATGCCAAT GAACTGCAAT TGGACGGTTT CCGTCTTGAT GCTGTCAAAC ACATTAAATT TTCTTTTTTG CGGGATTGGG TTAATCATGT GCAGTATTTT GAATGGTACA TGCCCAATGA CGGCCAACAT TGGAAGCGTT TGCAAAACGA CICGGCATAT TIGGCIGAAC ACGGIATIAC IGCCGICIGG AIICCCCCGG CATAIAAGGG AACGAGCCAA GCGGATGTGG GCTACGGTGC TTACGACCTT TATGATTTAG GGGAGTTTCA TCAAAAAGGG ACGGTTCGGA CAAAGTACGG CACAAAAGGA GAGCTGCAAT CTGCGATCAA AAGTCTTCAT TCCCGCGACA TTAACGTTTA CGGGGATGTG GTCATCAACC ACAAAGGCGG CGCTGATGCG ACCGAAGATG TAACCGCGGT TGAAGTCGAT CCCGCTGACC GCAACCGCGT AATITICAGGA GAACACCIAA ITAAAGCCIG GACACAITIT CAITITICCGG GGCGCGGCAG CACATACAGC GATTITAAAT GGCATTGGTA CCATTTTGAC GGAACCGATT GGGACGAGTC CCGAAAGCTG AACCGCATCT ATAAGTTTCA AGGAAAGGCT TGGGATTGGG AAGTTTCCAA TGAAAACGGC AACTATGATT ATTTGATGTA TGCCGACATC GATTATGACC ATCCTGATGT GCTCATCTTC TIGCTGCCTC ATTCTGCAGC AGCGGCGCA AATCTTAATG GGACGCTGAT

GAACGGTACG GTCGTTTCCA AGCATCCGTT GAAATCGGTT ACATTTGTCG ATAACCATGA SALI CGCGCTGGAA AACTATTTGA ACAAAACAAA TTTTAATCAT TCAGTGTTTG ACGTGCCGCT

ATGGCATGAC ATTACCGGAA ACCGTTCGGA GCCGGTTGTC ATCAATTCGG AAGGCTGGGG CGCTTTTATT CTCACAAGGG AATCTGGATA CCCTCAGGTT TTCTACGGGG ATATGTACGG AAAAGCGAGA AAACAGTATG CGTACGGAGC ACAGCATGAT TATTTCGACC ACCATGACAT TGTCGGCTGG ACAAGGGAAG GCGACAGCTC GGTTGCAAAT TCAGGTTTGG CGGCATTAAT AACAGACGGA CCCGGTGGGG CAAAGCGAAT GTATGTCGGC CGGCAAAACG CCGGTGAGAC AGAGTTTCAC GTAAACGGCG GGTCGGTTTC AATTTATGTT CAAAGATAGA AGAGCAGAGA GACGAAAGGA GACTCCCAGC GCGAAATTCC TGCCTTGAAA CACAAAATTG AACCGATCTT TACACAGCCG GGGCAATCGC TTGAGTCGAC TGTCCAAACA TGGTTTAAGC CGCTTGCTTA

BamHI

GITIGAATCG CATAGGTAAG GCGGGGATGA AATGGCAACG TTATCTGATG TAGCAAAGAA GGACGGATTT CCTGAAGGAA ATCCGTTTTT TTATTTTGCC CGTCTTATAA ATTTCTTTGA TTACATITITA TAATTIT TAACAAAGIG TCATCAGCCC TCAGGAAGGA CIIGCIGACA BclI

SEQ I'M NO: 5 AGCANATGTG TCGAAAATGA CGGTATCGCG GGTGATCA FIGURE 3

Oligonucleotide duplex A

SEQ ITS NO SEQ ITS NO			
NCO Bample Hindling NCO Bample Hindling	Oligonucleotide duplex B	$\frac{\text{Hga}_{\textbf{I}} \text{ SITE}}{\alpha - \text{AMYLASE}}$	5' CATG GCAAATCTTAATGGACGCTGATG 3' SEQ IN NO: S COTTTAGAATTACCTGCGACTACGTCAT 5' SEQ IN NO: 9 Met' mature α-Amylase